



ENTERED

PCT09

RAW SEQUENCE LISTING

DATE: 03/19/2003

PATENT APPLICATION: US/09/830,319B

TIME: 14:22:12

Input Set : A:\37921-151292.TXT

Output Set: N:\CRF4\03192003\I830319B.raw

4 <110> APPLICANT: BARTLETT, Perry F.
 5 HARTLEY, Lynne
 6 PILIZZOTTO, Mark
 7 KILPATRICK, Trevor
 8 KONTGEN, Frank
 9 COONAN, Jason
 10 GREFERATH, Ursula
 11 BOYD, Andrew W.
 12 GALEA, Mary
 13 PAXINOS, George
 14 MURPHY, Mark
 16 <120> TITLE OF INVENTION: A Method of Treatment
 19 <130> FILE REFERENCE: 37921-151292
 21 <140> CURRENT APPLICATION NUMBER: 09/830,319B
 22 <141> CURRENT FILING DATE: 2002-10-03
 24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
 25 <151> PRIOR FILING DATE: 1999-10-27
 27 <150> PRIOR APPLICATION NUMBER: PP6748
 28 <151> PRIOR FILING DATE: 1998-10-27
 30 <160> NUMBER OF SEQ ID NOS: 6
 32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 34 <210> SEQ ID NO: 1
 35 <211> LENGTH: 27
 36 <212> TYPE: DNA
 37 <213> ORGANISM: Artificial Sequence
 39 <220> FEATURE:
 40 <223> OTHER INFORMATION: PCR primer
 42 <400> SEQUENCE: 1
 43 cgtgctactt ccatttgtca cgtcctg
 45 <210> SEQ ID NO: 2
 46 <211> LENGTH: 21
 47 <212> TYPE: DNA
 48 <213> ORGANISM: Artificial Sequence
 50 <220> FEATURE:
 51 <223> OTHER INFORMATION: PCR primer
 53 <400> SEQUENCE: 2
 54 tgccgtgata gcaaatttga g
 56 <210> SEQ ID NO: 3
 57 <211> LENGTH: 22
 58 <212> TYPE: DNA
 59 <213> ORGANISM: Artificial Sequence
 61 <220> FEATURE:
 62 <223> OTHER INFORMATION: PCR primer

27

21

RAW SEQUENCE LISTING

DATE: 03/19/2003

PATENT APPLICATION: US/09/830,319B

TIME: 14:22:12

Input Set : A:\37921-151292.TXT

Output Set: N:\CRF4\03192003\I830319B.raw

```
64 <400> SEQUENCE: 3
65 aggaagtgag cattatggat ga 22
67 <210> SEQ ID NO: 4
68 <211> LENGTH: 20
69 <212> TYPE: DNA
70 <213> ORGANISM: Artificial Sequence
72 <220> FEATURE:
73 <223> OTHER INFORMATION: PCR primer
75 <400> SEQUENCE: 4
76 tgctcctcgt gccacgcgtt 20
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 27
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: PCR primer
86 <400> SEQUENCE: 5
87 ttagaattcc ccgaggagga gctgtac 27
89 <210> SEQ ID NO: 6
90 <211> LENGTH: 27
91 <212> TYPE: DNA
92 <213> ORGANISM: Artificial Sequence
94 <220> FEATURE:
95 <223> OTHER INFORMATION: PCR primer
97 <400> SEQUENCE: 6
98 ctagaattct gcagtccac caccocg 27
```

VERIFICATION SUMMARY

DATE: 03/19/2003

PATENT APPLICATION: US/09/830,319B

TIME: 14:22:13

Input Set : A:\37921-151292.TXT

Output Set: N:\CRF4\03192003\I830319B.raw

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/830,319A
Source: PT/09
Date Processed by STIC: 10/24/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/830,319A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length
Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
 (NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa; and which residue n or Xaa represents.
- 10 Invalid <213>
 Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



PCT09

RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:19

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

4 <110> APPLICANT: BARTLETT, Perry F.
 5 HARTLEY, Lynne
 6 PILIZZOTTO, Mark
 7 KILPATRICK, Trevor
 8 KONTGEN, Frank
 9 COONAN, Jason
 10 GREFERATH, Ursula
 11 BOYD, Andrew W.
 12 GALEA, Mary
 13 PAXINOS, George
 14 MURPHY, Mark
 16 <120> TITLE OF INVENTION: A Method of Treatment
 19 <130> FILE REFERENCE: 37921-151292
 21 <140> CURRENT APPLICATION NUMBER: 09/830,319A
 22 <141> CURRENT FILING DATE: 2002-10-03
 24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
 25 <151> PRIOR FILING DATE: 1999-10-27
 27 <150> PRIOR APPLICATION NUMBER: PP6748
 28 <151> PRIOR FILING DATE: 1998-10-27
 30 <160> NUMBER OF SEQ ID NOS: 6
 32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 34 <210> SEQ ID NO: 1
 35 <211> LENGTH: 27
 36 <212> TYPE: DNA
 37 <213> ORGANISM: Oligonucleotides
 39 <400> SEQUENCE: 1
 40 cgtgctactt ccatttgtca cgtcctg
 42 <210> SEQ ID NO: 2
 43 <211> LENGTH: 21
 44 <212> TYPE: DNA
 45 <213> ORGANISM: Oligonucleotides
 47 <400> SEQUENCE: 2
 48 tgccgtgata gcaaatttga g
 50 <210> SEQ ID NO: 3
 51 <211> LENGTH: 22
 52 <212> TYPE: DNA
 53 <213> ORGANISM: Oligonucleotides
 55 <400> SEQUENCE: 3
 56 aggaagttag cattatggat ga
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 20
 60 <212> TYPE: DNA
 61 <213> ORGANISM: Oligonucleotides

Does Not Comply
Corrected Diskette Needed

pg 1-2

OK->

(global error)
invalid response - see item 10 on Error
summary
Sheet

27

21

22

RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:19

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

63 <400> SEQUENCE: 4
64 tgctcctcgt gccagcggt 20
66 <210> SEQ ID NO: 5
67 <211> LENGTH: 27
68 <212> TYPE: DNA
69 <213> ORGANISM: Oligonucleotides
71 <400> SEQUENCE: 5
72 ttagaattcc ccgaggagga gctgtac 27
74 <210> SEQ ID NO: 6
75 <211> LENGTH: 27
76 <212> TYPE: DNA
77 <213> ORGANISM: Oligonucleotides
79 <400> SEQUENCE: 6
80 ctagaattct gcagtccac caccgcg 27

VERIFICATION SUMMARY

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:20

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date

FY



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/830,319A
Source: pc/09
Date Processed by STIC: 10/24/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/830,319A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length
Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES)
Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
 (NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220>
Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.



PCT09

RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:19

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

4 <110> APPLICANT: BARTLETT, Perry F.
 5 HARTLEY, Lynne
 6 PILIZZOTTO, Mark
 7 KILPATRICK, Trevor
 8 KONTGEN, Frank
 9 COONAN, Jason
 10 GREFERATH, Ursula
 11 BOYD, Andrew W.
 12 GALEA, Mary
 13 PAXINOS, George
 14 MURPHY, Mark
 16 <120> TITLE OF INVENTION: A Method of Treatment
 19 <130> FILE REFERENCE: 37921-151292
 21 <140> CURRENT APPLICATION NUMBER: 09/830,319A
 22 <141> CURRENT FILING DATE: 2002-10-03
 24 <150> PRIOR APPLICATION NUMBER: PCT/AU99/00931
 25 <151> PRIOR FILING DATE: 1999-10-27
 27 <150> PRIOR APPLICATION NUMBER: PP6748
 28 <151> PRIOR FILING DATE: 1998-10-27
 30 <160> NUMBER OF SEQ ID NOS: 6
 32 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 34 <210> SEQ ID NO: 1
 35 <211> LENGTH: 27
 36 <212> TYPE: DNA
 37 <213> ORGANISM: Oligonucleotides
 39 <400> SEQUENCE: 1
 40 cgtgctactt ccatttgtca cgtcctg
 42 <210> SEQ ID NO: 2
 43 <211> LENGTH: 21
 44 <212> TYPE: DNA
 45 <213> ORGANISM: Oligonucleotides
 47 <400> SEQUENCE: 2
 48 tgccgtgata gcaaatttga g
 50 <210> SEQ ID NO: 3
 51 <211> LENGTH: 22
 52 <212> TYPE: DNA
 53 <213> ORGANISM: Oligonucleotides
 55 <400> SEQUENCE: 3
 56 aggaagtga cattatggat ga
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 20
 60 <212> TYPE: DNA
 61 <213> ORGANISM: Oligonucleotides

Does Not Comply
Corrected Diskette Needed

pg 1-2

OK->

(global error)
invalid response - see item 10 on End
summary
sheet

27

21

22

RAW SEQUENCE LISTING

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:19

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

63 <400> SEQUENCE: 4
64 tgctcctcgt gccagcggt 20
66 <210> SEQ ID NO: 5
67 <211> LENGTH: 27
68 <212> TYPE: DNA
69 <213> ORGANISM: Oligonucleotides
71 <400> SEQUENCE: 5
72 ttagaattcc ccgaggagga gctgtac 27
74 <210> SEQ ID NO: 6
75 <211> LENGTH: 27
76 <212> TYPE: DNA
77 <213> ORGANISM: Oligonucleotides
79 <400> SEQUENCE: 6
80 ctagaattct gcagtccac caccg 27

VERIFICATION SUMMARY

DATE: 10/24/2002

PATENT APPLICATION: US/09/830,319A

TIME: 13:42:20

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\10242002\I830319A.raw

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

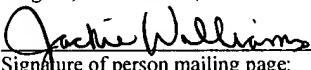
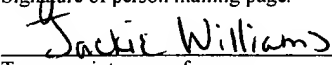
In re:	Patent application of Perry F. Bartlett, et al.	:	
		:	
		:	Group Art Unit:
Serial No.	09/830,319	:	
		:	
Filing Date:	(Int'l.)October 27, 1999	:	Examiner:
		:	
For:	A Method of Treatment	:	

STATEMENT PURSUANT TO 37 CFR § 1.825

Commissioner for Patents
Box PCT
Washington, D.C. 20231

Sir:

The Substitute Sequence Listing filed herewith includes no new matter. The content of the substitute Sequence Listing in computer readable form is the same as the substitute Sequence Listing paper copy submitted herewith.

CERTIFICATE OF MAILING UNDER 37 C.F.R. 1.10
EXPRESS MAIL Mailing Label Number: <u>EL931 089 170 US</u>
Date of Deposit: <u>11/21/02</u>
<p>I hereby certify that this correspondence, along with any paper referred to as being attached or enclosed, and/or fee, is being deposited with the United States Postal Service, "EXPRESS MAIL-POST OFFICE TO ADDRESSEE" service under 37 CFR 1.10, on the date indicated above, and addressed to: Commissioner for Patents, Washington, D.C. 20231.</p>
 _____ Signature of person mailing page:
 _____ Type or print name of person

Respectfully submitted,

PERRY F. BARTLETT, et al

BY 

DANIEL A. MONACO
Reg. No. 30,480
DRINKER, BIDDLE & REATH, LLP.
One Logan Square
18th and Cherry Streets
Philadelphia, PA 19103-6996
(215) 988-3312
(215) 988-2757 – fax
Attorney for the Applicants